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Taxonomic analysis of 16S rRNA reads extracted from metagenome

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An abbreviated version of this protocol was published in Science in Nov 2019

Pathogen-induced activation of disease-suppressive functions in the endophytic root microbiome

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How to cite:(Readers should cite both the Bio-protocol preprint and the original research article where this protocol was used)

1. Carrión, V. , Cordovez, V. , Medema, M. and Raaijmakers, J. (2023). Taxonomic analysis of 16S rRNA reads extracted from metagenome. Bio-protocol Preprint. bio-protocol.org/prep2115.
2. Carrión, V. J., Perez-Jaramillo, J., Cordovez, V., Tracanna, V., Hollander, M. D., Ruiz-Buck, D., Mendes, L. W., Ijcken, W. F. V., Gomez-Exposito, R., Elsayed, S. S., Mohanraju, P., Arifah, A., Oost, J. V. D., Paulson, J. N., Mendes, R., Wezel, G. P. V., Medema, M. H. and Raaijmakers, J. M. (2019). Pathogen-induced activation of disease-suppressive functions in the endophytic root microbiome. Science 366(6465). DOI: 10.1126/science.aaw9285

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